



Gene ID	Description	Number of unique peptides	PMSS	% Coverage
Tb927.2.4710	RNA-binding protein (RRM1)	8	105.7	29
Tb927.11.14100	PTB2 (DRBD4)	2	13.3	5
Tb927.6.2550	RNA-binding protein (RRM2)	2	13.1	4
Tb05.5K5.70 Tb927.5.4420	Nucleolar RNA helicase II, putative, nucleolar RNA helicase Gu	4	36	9
Tb927.2.100	Retrotransposon hot spot protein 1 (RHS1)	2	13.2	3
Tb927.2.1120	Retrotransposon hot spot protein 4 (RHS4)	7	70.8	8
Tb927.2.1080	Retrotransposon hot spot protein 5 (RHS5)	2	12.2	3
Tb927.1.2370	Beta tubulin	5	35.2	22
Tb927.1.2380	Alpha tubulin	2	14.5	8
Tb927.10.6400	Chaperonin Hsp60, mitochondrial precursor	3	21.4	8
Tb927.6.3800	Heat shock 70 kDa protein, mitochondrial precursor	3	21	8
Tb927.11.11330	Heat shock protein 70	3	18.4	6
Tb927.10.10900	Heat shock protein, putative HSP90	2	13.6	4
Tb927.7.2910	Histone H2A	2	48	32
Tb927.10.10540	Histone H2B	3	58.2	41
Tb927.1.2450	Histone H3	4	27.7	32
Tb927.2.2670	Histone H4	6	125.4	37

Figure S2. TbRRM1 interacting proteins.

Immunoprecipitation of HA-RRM1 (lane1) and wild-type (lane2) parasite lysates was performed with anti-HA matrix and proteins were separated by SDS-PAGE. Bands subjected to LC-MS are indicated by arrows. Bands present in both lanes were not selected for MS identification. Antibody light chain is indicated by an asterisk. The table shows HA-RRM1 interacting proteins from selected bands